(FILE 'HOME' ENTERED AT 10:44:14 ON 28 JAN 2003)

FILE 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH, AGRICOLA' ENTERED AT

10:44:36 ON 28 JAN 2003

- L1 10512953 S PEPTIDE OR POLYPEPTIDE OR OLIGOPEPTIDE OR PROTEIN OR RECEPTOR
- L2 13965 S (FACTOR VIIA) OR (FACTOR VIIAI) OR (FFR-FACTOR VIIA) OR (GLUC
- L3 1457073 S (L1 OR L2) (P) (PURIF? OR ISOLAT?)
- L4 30144 S ANION (W) EXCHANGE (W) (CHROMATOGRAPHY OR COLUMN)
- L5 8956 S L3 (P) L4
- L6 5826 S ORGANIC MODIFIER
- L7 890678 S ALKANOL OR ALKYNOL OR ALKENOL OR UREA OR GUANIDINE OR (ALKNAO
- L8 0 S L5 (P) L6
- L9 527 S L5 (P) L7
- L10 0 S L9 (P) IMPURIT?
- L11 103 S L9 (P) ELUT?
- L12 37 DUPLICATE REMOVE L11 (66 DUPLICATES REMOVED)
- L13 0 S L9 (P) (NEGATIVE CHARGE)

 $=> \log y$

		2003/01/2 8 09:47	DERWENT	C1-6-alkanol	Ц	L7	BRS	7
i		2003/01/2	USPAT; US-PGPUB; EPO; JPO; DERWENT	organic adj modifier	374	L6	BRS	9
		2003/01/2 8 09:45	USPAT; US-PGPUB; EPO; JPO; DERWENT	3 same 4	1643	L5	BRS	Л
		2003/01/2 8 09:45	USPAT; US-PGPUB; EPO; JPO; DERWENT	anion adj exchange adj (chromatography or column)	5161	L4	BRS	4
		2003/01/2 8 09:44	USPAT; US-PGPUB; EPO; JPO; DERWENT	1 same 2	10506 2	Ь3	BRS	ω
		2003/01/2 8 09:44	USPAT; US-PGPUB; EPO; JPO; DERWENT	purif\$7 or isolat\$3	10404 69	L2	BRS	2
		2003/01/2	USPAT; US-PGPUB; EPO; JPO; DERWENT	peptide or polypeptide or oligopeptide or protein or receptor or vira or glucagon or hgh orinsulin or (factor adj VIIa) or (factor adj viiai) or (factor adj viiai) or (FFR-factor adj VIIa) or (glucagon-like adj peptide-1) or (glucagon-like adj peptide-2)	41219 4	L1	BRS	P
Err or Def ini tio	Comm	Time Stamp	DBs	Search Text	Hits	#	Туре	

0		2003/01/2 8 10:01	USPAT; US-PGPUB; EPO; JPO; DERWENT	staby adj arne.in.	2	L15	BRS	15
0		2003/01/2 8 10:01	USPAT; US-PGPUB; EPO; JPO; DERWENT	9 same 13	2	L14	BRS	14
0		2003/01/2 8 09:56 ·	USPAT; US-PGPUB; ; EPO; JPO; DERWENT	industrial	45875 9	L13	BRS	13
0		2003/01/2 8 09:56	USPAT; US-PGPUB; EPO; JPO; DERWENT	9 same impurity	4	L12	BRS	12
0		2003/01/2 8 09:52	PGPUB;	9 same 10	- -	L11	BRS	11
0		2003/01/2 8 09:50		28079negative adj charge	28079	L10	BRS	10
0		2003/01/2 8 09:49	USPAT; US-PGPUB; EPO; JPO; DERWENT	5 same (6 or 8)	85	Г9	BRS	9
0		2003/01/2 8 09:49	USPAT; US-PGPUB; EPO; JPO; DERWENT	alkanol or alkylnol or alkenol or urea or guanidine or (alkanoic adj acid) or polyalcohol	22676	18	BRS	ω
# H H	Err or Def ts ini tio	Time Comm Stamp ents	DBs	Search Text	Hits	#	Туре	

*

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s alkanol or alkynol or alkenol or urea or guanidine or (alknaoic acid) or glycol or polyalcoho

890678 ALKANOL OR ALKYNOL OR ALKENOL OR UREA OR GUANIDINE OR (ALKNAOIC

s organic modifier

5826 ORGANIC MODIFIER

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> s 15 (p) 16
            0 L5 (P) L6
> s 15 (p) 17
          527 L5 (P) L7
> s 19 (p) impurit?
            0 L9 (P) IMPURIT?
s 19 (p) elut?
          103 L9 (P) ELUT?
duplicate remove 111
JPLICATE PREFERENCE IS 'MEDLINE, CAPLUS, BIOSIS, EMBASE, SCISEARCH, AGRICOLA'
EEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n
ROCESSING COMPLETED FOR L11
            37 DUPLICATE REMOVE L11 (66 DUPLICATES REMOVED)
s 19 (p) (negative charge)
            0 L9 (P) (NEGATIVE CHARGE)
> d 112 1-37 ibib abs
L2 ANSWER 1 OF 37 CAPLUS COPYRIGHT 2003 ACS
CCESSION NUMBER: 2002:637703 CAPLUS
CUMENT NUMBER:
                          137:181920
                          Method for purifying the Helicobacter adhesin-like
TLE:
                          protein A (AlpA)
                          Fourrichon, Laurence; Lissolo, Ling; Pitiot, Olivier
IVENTOR(S):
ATENT ASSIGNEE(S):
                       Merieux Oravax, Fr.
OURCE:
                          PCT Int. Appl., 22 pp.
                          CODEN: PIXXD2
CUMENT TYPE:
                          Patent
ANGUAGE:
                          French
AMILY ACC. NUM. COUNT: 1
ATENT INFORMATION:
                             DATE APPLICATION NO. DATE
   PATENT NO. KIND DATE
                      ----
   WO 2002064622 A1 20020822 WO 2002-FR355 20020130
       W: AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH,
            PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZM, ZW, AM, AZ, BY, KG, KZ, MD, RU,
            TJ, TM
       RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG
                       A1 20020809
                                             FR 2001-1499
   FR 2820424
                                                                  20010205
RIORITY APPLN. INFO.:
   THE invention concerns a method for ***purifying*** the Helicobacter
                                           FR 2001-1499
   adhesin-like ***protein*** A (AlpA) which consists in: (i) contacting
   an AlpA prepn. and 2.5 to 3.5 M of ***guanidine*** with a hydrophobic
   interaction chromatog. material, so that the AlpA is adsorbed on the
                          ***eluting*** the AlpA with a soln. contg. 3.5 to
   material; and (ii)
   4.5 M of ***guanidine*** . The AlpA prepn. to be ***purified***
   can be in particular derived from an E. coli culture capable of expressing
   AlpA in a high-level recombinant form, rAlpA being in the form of
   inclusion bodies, the latter being recovered and solubilized in the
   presence of ***guanidine*** , and optionally ammonium sulfate-pptd. for
   the purpose of preliminary ***purifn*** . The hydrophobic interaction
   chromatog. can be followed up by an ***anion*** ***exchange***
     ***chromatog*** . in the presence of 8 M of ***urea***
FERENCE COUNT:
                                THERE ARE 7 CITED REFERENCES AVAILABLE FOR THIS
                                 RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT
```

2 ANSWER 2 OF 37

CESSION NUMBER: 2000290831

MEDLINE

MEDLINE

DUPLICATE 1

CUMENT NUMBER:

LE:

20290831 PubMed ID: 10828960 Monoclonal antilly light chain with prothrombin

activity.

Thiagarajan P; Dannenbring R; Matsuura K; Tramontano A; THOR:

Gololobov G; Paul S

Departments of Internal Medicine and Pathology and RPORATE SOURCE:

Laboratory Medicine, Center for Chemical Immunology, University of Texas-Houston Medical School, 77030, USA...

Perumal.Thiagarajan@uth.tmc.edu

AI 31268 (NIAID) NTRACT NUMBER:

HL 44126 (NHLBI) HL 65096 (NHLBI)

BIOCHEMISTRY, (2000 May 30) 39 (21) 6459-65. JRCE:

Journal code: 0370623. ISSN: 0006-2960.

United States 3. COUNTRY:

Journal; Article; (JOURNAL ARTICLE) CUMENT TYPE:

English **IGUAGE:**

Priority Journals LE SEGMENT:

200007 TRY MONTH:

Entered STN: 20000720 TRY DATE:

Last Updated on STN: 20000720

Entered Medline: 20000711

Prothrombin is the precursor of thrombin, a central enzyme in coagulation. Autoantibodies to prothrombin are associated with thromboembolism, but the mechanisms by which the antibodies modulate the coagulation processes are not understood. We screened a panel of 34 monoclonal antibody light chains from patients with multiple myeloma for prothrombinase ***isolated*** activity by an electrophoresis method. Two light chains with the activity were identified, and one of the light chains was characterized further. The prothrombinase activity ***eluted*** from a gel-filtration column run in denaturing solvent (6 M ***guanidine*** hydrochloride) at the characteristic positions of the light chain dimer and monomer. A constant level of catalytic activity was observed across the width of the light chain monomer peak, assessed as the cleavage of IEGR-methylcoumarinamide, substrate corresponding to residues 268-271 of ***peptide*** ***peptide*** by the light chain was prothrombin. Hydrolysis of this saturable and consistent with Michaelis-Menten-Henri kinetics (K(m) 103 microM; k(cat) of 2.62 x 10(-)(2)/min). Four cleavage sites in prothrombin were identified by N-terminal sequencing of the fragments: Arg(155)-Ser(156), Arg(271)-Thr(272), Arg(284)-Thr(285), and Arg(393)-Ser(394). The light chain did not cleave radiolabeled albumin, thyroglobulin, and annexin V under conditions that readily permitted detectable prothrombin cleavage. Two prothrombin fragments (M(r) 55 000 ***anion*** - ***exchange*** ***isolated*** by and 38 000), were ***chromatography*** and were observed to cleave a thrombin substrate, tosyl-GPR-nitroanilide. Conversion of fibrinogen to fibrin was accelerated by the prothrombin fragments generated by the light chain. These finding suggest a novel mechanism whereby antibodies can induce a procoagulant state, i.e., prothrombin activation via cleavage of the molecule.

DUPLICATE 2 MEDLINE 2 ANSWER 3 OF 37

2002257174 MEDLINE CESSION NUMBER:

PubMed ID: 11996098 21991123 CUMENT NUMBER:

Proteoglycans of human umbilical cord arteries. TLE:

Gogiel T; Jaworski S THOR: RPORATE SOURCE:

Department of Biochemistry, Medical Academy of Bialystok,

Poland.. tgogiel@amb.ac.bialystok.pl

ACTA BIOCHIMICA POLONICA, (2000) 47 (4) 1081-91.

Journal code: 14520300R. ISSN: 0001-527X.

B. COUNTRY: Poland

URCE:

Journal; Article; (JOURNAL ARTICLE) CUMENT TYPE:

English MGUAGE:

LE SEGMENT: Priority Journals

TRY MONTH: 200206

Entered STN: 20020509 ITRY DATE:

Last Updated on STN: 20020625 Entered Medline: 20020624

Proteoglycans (PGs) were dissociatively extracted from human umbilical cord arteries (UCAs) with 4 M ***guanidine*** hydrochloride containing Triton X-100 and protease inhibitors, ***purified*** by Q-Sepharose analyzed by gel filtration SDS/PAGE and agarose gel

electrophoresis before and aft treatment with chondroitinase found that the PG preparation especially enriched in chondroitin/dermatan sulphate PGs. The predominant PG fraction included small PGs that emerged from Sepharose CL-2B with Kav = 0.74. Their molecular mass, estimated by SDS/PAGE, was 160-200 kDa and 90-150 kDa, i.e. it was typical for biglycan and decorin, respectively. Treatment with chondroitinase ABC yielded the core ***proteins*** of 45 and 47 kDa, characteristic for both small PGs. Remarkable amounts of the 45 kDa ***protein*** were detected in non-treated PG samples, suggesting the presence of free core ***proteins*** of biglycan and decorin. Large PGs were present in lower amounts. In intact form they were from Sepharose CL-2B with Kav = 0.17 and 0.43. Digestion with ***proteins*** with a molecular chondroitinase ABC yielded the core mass within the range of 180-360 kDa but predominant were the bands of 200, 250 and 360 kDa. The large PGs probably represent various forms of versican or perlecan bearing chondroitin sulphate chains.

DUPLICATE 3 2 ANSWER 4 OF 37 MEDLINE CESSION NUMBER: 2001434257 MEDLINE 21094863 PubMed ID: 11162736 CUMENT NUMBER: A new purification method for overproduced proteins $\mathtt{TLE}:$ sensitive to endogenous proteases. Saijo-Hamano Y; Namba K; Oosawa K THOR:

Protonic NanoMachine Project, ERATO, JST, 1-7 Hikaridai,

Seika 619-0237, Japan.

JOURNAL OF STRUCTURAL BIOLOGY, (2000 Nov) 132 (2) 142-6.

Journal code: 9011206. ISSN: 1047-8477.

United States B. COUNTRY:

Journal; Article; (JOURNAL ARTICLE) CUMENT TYPE:

English NGUAGE:

RPORATE SOURCE:

URCE:

Priority Journals LE SEGMENT:

200108 TRY MONTH:

Entered STN: 20010806 TRY DATE:

Last Updated on STN: 20010806 Entered Medline: 20010802

Proteolysisis a major problem in ***purification*** of overproduced ***proteins*** for structural studies. We developed a new method to avoid proteolysis of the products even in cases where popular protease inhibitors do not work effectively. When we cloned FlgF, a flagellar rod

protein , from Salmonella typhimurium and overproduced it in Escherichia coli, FlgF was highly susceptible to cleavage by endogenous proteases after cell disruption even in the presence of various protease inhibitors. However, FlgF was not digested when the cells were disrupted in the presence of ***urea*** , which allowed us to develop the following new ***purification*** procedure. After cell disruption in the presence of ***urea*** and removal of the cell debris, the supernatant was passed through tandem-connected cation- and cation-exchange column, and protease-free FlgF was ***eluted*** from ***column*** . This the disconnected ***anion*** - ***exchange*** gave a stable full-length product suitable for crystallization trials. The key procedures are cell disruption in the presence of ***urea*** linked ion-exchange chromatography to quickly remove proteases as well as ***urea*** . This fast and simple method can be applied to

purification of other overproduced ***proteins*** very sensitive to proteolysis.

Copyright 2000 Academic Press.

2 ANSWER 5 OF 37 CAPLUS COPYRIGHT 2003 ACS 1999:235191 CAPLUS CESSION NUMBER:

131:41348 CUMENT NUMBER:

Purification and properties of a cholesteryl ester TLE:

hydrolase from rat liver microsomes

Cristobal, Susana; Ochoa, Begona; Fresnedo, Olatz JTHOR(S): RPORATE SOURCE:

Department of Physiology, University of the Basque Country Medical School, Bilbao, 48080, Spain Journal of Lipid Research (1999), 40(4), 715-725

CODEN: JLPRAW; ISSN: 0022-2275

Lipid Research, Inc. JBLISHER:

CUMENT TYPE: Journal

OURCE:

ANGUAGE: English 1 actor budrolace (CEH) from

female rat liver microsomes, ar some structural, immunol., kin regulatory properties of the eleme that distinguish microsomal other hepatic cholesteryl ester-splitting enzymes are reported. CEH was 12.4-fold from re- ***isolated*** microsomes using ***purified*** sequential solubilization by sonication, polyethylene ***glycol*** pptn., fractionation with hydroxylapatite, ***anion*** ***chromatog*** ., and chromatog. on hydroxylapatite, ***exchange*** with an overall yield of 3.2%. CEH activity was ***purified*** 141-fold over nonspecific esterase activity and 56-fold over triacylglycerol lipase activity. In sharp contrast to most esterases and lipases, CEH did not bind to Con A-Sepharose and heparin-Sepharose. After PAGE, the ***purified*** enzyme exhibited 2 Ag-stained bands, but only the ***protein*** electro- ***eluted*** from the low-mobility band had CEH activity. Affinity- ***purified*** polyclonal antibodies raised to electro- ***eluted*** CEH inhibited 90% of the activity of liver microsomal CEH and reacted with a 106-kDa ***protein*** Western blot anal. This 106-kDa CEH contained a unique N-terminal amino acid sequence. The ***purified*** enzyme had optimal activity at pH 6 and no taurocholate requirement, and was inhibited by the serine active site inhibitor, phenylmethylsulfonyl fluoride, and by free SH group-specific reagents. It hydrolyzed cholesteryl oleate much more efficiently than triolein, and its hydrolytic activity with p-nitrophenyl acetate was higher than with p-nitrophenyl butyrate. Thus, the results indicate that rat liver microsomes contain a bile salt-independent ***protein*** that is relatively specific for cholesteryl catalytic ester hydrolysis. THERE ARE 48 CITED REFERENCES AVAILABLE FOR THIS 48 FERENCE COUNT: RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT DUPLICATE 4 ANSWER 6 OF 37 MEDLINE CESSION NUMBER: 1999133646 MEDLINE CUMENT NUMBER: 99133646 PubMed ID: 9950147 Purification and characterization of a complex from TLE: placental syncytiotrophoblast microvillous membranes which inhibits the proliferation of human umbilical vein endothelial cells. Kertesz Z; Hurst G; Ward M; Willis A C; Caro H; Linton E A; THOR: Sargent I L; Redman C W Nuffield Department of Obstetrics and Gynaecology, RPORATE SOURCE: University of Oxford, UK.. zkertesz@radius.jr2.ox.ac.uk PLACENTA, (1999 Jan) 20 (1) 71-9. URCE: Journal code: 8006349. ISSN: 0143-4004. B. COUNTRY: ENGLAND: United Kingdom Journal; Article; (JOURNAL ARTICLE) English NGUAGE: LE SEGMENT: Priority Journals TRY MONTH: 199903

CUMENT TYPE:

TRY DATE: Entered STN: 19990413

> Last Updated on STN: 20021030 Entered Medline: 19990329

The signs of pre-eclampsia are thought to arise from maternal endothelial dysfunction caused by circulating factors of placental origin. Syncytiotrophoblast microvillous membranes (STBM) cause endothelial disruption and inhibit proliferation in vitro. Significantly increased amounts of STBM can be detected in blood from pre-eclamptic women and could contribute to endothelial dysfunction in vivo. This study

purified a complex from STBM which inhibits the proliferation of cultured human endothelial cells. Integral membrane ***proteins*** were solubilized with sucrose monolaurate. ***Anion***

yielded two peaks of ***exchange*** ***chromatography*** anti-proliferative activity. Only the second peak was specific to STBM and was subjected to further separation by Sephacryl S-200 gel filtration chromatography (GFC). A single peak of specific activity ***eluted*** close to the void volume, at a position unaltered by added denaturing agents, guanidium chloride or ***urea*** . On Sephacryl S-300 GFC, two peaks were obtained of 410 and 820 kDa, with similar anti-proliferative ***protein*** components (by SDS-polyacrylamide gel activity and electrophoresis). The major ***protein*** bands were as integrins alpha5 and alpha v, dipeptidyl peptidase IV, alpha-actinin, transferrin, transferrin ***receptor*** , placental alkaline phosphatase and monoamine oxidase A.

DUPLICATE 5 ANSWER 7 OF 37 MEDLINE

1998334669 SSION NUMBER:

PubMed ID: 9668118 98334669 MENT NUMBER:

Identification and characterization of a bovine neurite Æ:

growth inhibitor (bNI-220).

Spillmann A A; Bandtlow C E; Lottspeich F; Keller F; Schwab IOR:

Brain Research Institute, University of Zurich and Swiss PORATE SOURCE:

Federal Institute of Technology Zurich, August Forelstrasse

1, 8029 Zurich, Switzerland.

JOURNAL OF BIOLOGICAL CHEMISTRY, (1998 Jul 24) 273 (30) RCE:

19283-93.

Journal code: 2985121R. ISSN: 0021-9258.

United States . COUNTRY:

Journal; Article; (JOURNAL ARTICLE) UMENT TYPE:

GUAGE: English

Priority Journals E SEGMENT:

199808 RY MONTH:

RY DATE: Entered STN: 19980828

Last Updated on STN: 19980828 Entered Medline: 19980820

The poor axonal regeneration that follows lesions of the central nervous system (CNS) is crucially influenced by the local CNS tissue environment through which neurites have to grow. In addition to an inhibitory role of the glial scar, inhibitory substrate effects of CNS myelin and ***proteins*** oligodendrocytes have been demonstrated. Several including NI-35/250, myelin-associated glycoprotein, tenascin-R, and NG-2 have been described to have neurite outgrowth inhibitory or repulsive properties in vitro. Antibodies raised against NI-35/250 (monoclonal antibody IN-1) were shown to partially neutralize the growth inhibitory effect of CNS myelin and oligodendrocytes, and to result in long distance fiber regeneration in the lesioned adult mammalian CNS in vivo. We report ***purification*** of a myelin ***protein*** to apparent here the homogeneity from bovine spinal cord which exerts a potent neurite outgrowth inhibitory effect on PC12 cells and chick dorsal root ganglion cells, induces collapse of growth cones of chick dorsal root ganglion cells, and also inhibits the spreading of 3T3 fibroblasts. These activities could be neutralized by the monoclonal antibody IN-1. The ***purification*** procedure includes detergent solubilization, ***anion*** ***elution*** from high resolution SDS-polyacrylamide gel electrophoresis. The active ***protein*** has a molecular mass of 220

kDa and an isoelectric point between 5.9 and 6.2. Its inhibitory activity is sensitive to protease treatment and resists harsh treatments like 9 M ***urea*** or short heating. Glycosylation is, if present at all, not detectable. Microsequencing resulted in six ***peptides*** strongly suggests that this ***proteins*** is novel.

2 ANSWER 8 OF 37 CAPLUS COPYRIGHT 2003 ACS 1997:416857 CAPLUS CESSION NUMBER:

CUMENT NUMBER: 127:39795

Anion exchange process for the purification of Factor TLE:

Bhattacharya, Prabir; Motokubota, Toshiharu; Fedalizo, IVENTOR(S):

Norman M.

Alpha Therapeutic Corporation, USA ATENT ASSIGNEE(S):

PCT Int. Appl., 29 pp.

CODEN: PIXXD2

CUMENT TYPE: Patent English ANGUAGE:

AMILY ACC. NUM. COUNT:

ATENT INFORMATION:

OURCE:

PATENT NO. KIND DATE APPLICATION NO. DATE ______ ______ WO 9717370 WO 1996-US17806 19961107 A1 19970515

W: CA, JP

RW: AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE A 19970819 US 1995-554724 : 19951107 US 5659017 US 1995-554724 19951107 RIORITY APPLN. INFO.:

A new method for ***purifying*** Factor VIII complex from an impure " " a The arvannt dissolved in a heparin soln. In Factor VIII complex is initially ***purified*** by polyethy the ***glycol*** pptn. The tor VIII contg. supernatant collected after the pptn. is loaded into an ***anion*** ***exchange*** ***column*** that has a quaternary amino Et group. The Factor VIII complex is then ***eluted*** from the column with a buffer comprising from about 0.14M to about 0.20M CaCl2. The final step in the ***purifn*** . is to ppt. the Factor VIII complex in the presence of glycine and sodium chloride. The pptd. Factor VIII complex is then reconstituted and stabilized. The reconstituted Factor VIII complex can then be lyophilized and dry heated to obtain a final Factor VIII product.

2 ANSWER 9 OF 37 CAPLUS COPYRIGHT 2003 ACS CESSION NUMBER: 1997:695207 CAPLUS

CUMENT NUMBER: 127:344221

TLE: Characteristics of chondroitin sulfated proteoglycans

in the matrix phase of human alveolar bone

THOR(S): Zhao, Hu; Suzuki, Naoto; Maeno, Masao; Katayama,

Ichiro; Arai, Toshiyuki; Otsuka, Kichibee

RPORATE SOURCE: Department of Biochemistry, Nihon University School of

Dentistry, Kanda-surugadai, 101, Japan

Journal of Hard Tissue Biology (1997), 6(1), 1-9

CODEN: JHTBFF; ISSN: 1341-7649

BLISHER: Japanese Society of Hard Tissue Research & Technology

CUMENT TYPE: Journal NGUAGE: English

URCE:

This study demonstrates the types and characteristics of chondroitin sulfated proteoglycans (CSPGs) in the matrix phase of human alveolar bone (HAB) extd. with ***guanidine*** -HCl from EDTA-demineralized HAB fragments. Different CSPGs including dermatan sulfated PG were analyzed and identified using three monoclonal antibodies in combination with different chondroitinase digestions. The PGs were partially

purified by gel filtration and then ***anion*** ***exchange*** ***column*** chromatog. under denaturing conditions.
The CSPGs in each ***eluted*** fraction from the columns were detected
by ELISA using a monoclonal antibody (MAb) 2B6 in combination with
chondroitinase ABC digestion. 2B6-Reactive fractions of the ***anion***
- ***exchange*** ***column*** were concd. in three fractions. The
types of glycosaminoglycan (GAG) chains of CSPGs were detd. by Western
blotting using antibodies against GAG stubs on a core ***protein***
after enzyme digestion. The main CSPGs in the matrix phase of HAB were
comprised of dermatan sulfated PG contg. a 45 kDa core ***protein***.
There were also small amts. of chondroitin non-sulfated PGs which
consisted of 45 and 110 kDa core ***proteins***, but no chondroitin

.2 ANSWER 10 OF 37 MEDLINE DUPLICATE 6

CESSION NUMBER: 95357413 MEDLINE

6-sulfated PG was detected.

CUMENT NUMBER: 95357413 PubMed ID: 7630943

TLE: Purification of NAD-dependent mannitol dehydrogenase from

celery suspension cultures.

JTHOR: Stoop J M; Williamson J D; Conkling M A; Pharr D M

)RPORATE SOURCE: Department of Horticultural Science, North Carolina State

University, Raleigh 27695-7609, USA.

DURCE: PLANT PHYSIOLOGY, (1995 Jul) 108 (3) 1219-25.

Journal code: 0401224. ISSN: 0032-0889.

JB. COUNTRY: United States

)CUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

ANGUAGE: English

[LE SEGMENT: Priority Journals

ITRY MONTH: 199509

NTRY DATE: Entered STN: 19950921

Last Updated on STN: 20020420 Entered Medline: 19950905

Mannitol dehydrogenase, a mannitol:mannose 1-oxidoreductase, constitutes the first enzymatic step in the catabolism of mannitol in nonphotosynthetic tissues of celery (Apium graveolens L.). Endogenous regulation on the enzyme activity in response to environmental cues is critical in modulating tissue concentration of mannitol, which, importantly, contribute to stress tolerance of celery. The enzyme was ***purified*** to homogeneity from celery suspension cultures grown on

```
***purified*** 589-fold to specific activity of 365 mumol 1 mg-1
***protein*** with a 37% y d of enzyme activity present in the crude
    .***purified***
   extract. A highly efficient and simple ***purification***
                                                                   protocol was
   developed involving polyethylene ***glycol***
diethylaminoethyl- ***anion*** - ***exchange***
                                                     fractionation,
     ***chromatography*** , and NAD-agarose affinity chromatography using NAD
             ***elution*** . Sodium dodecylsulfate gel electrophoresis of
   the final preparation revealed a single 40-kD ***protein*** . The
                                 ***protein***
                                                  was determined to be
   molecular mass of the native
   approximately 43 kD, indicating that the enzyme is a monomer. Polyclonal
   antibodies raised against the enzyme inhibited enzymatic activity of
                     mannitol dehydrogenase. Immunoblots of crude
     ***purified***
                     extracts from mannitol-grown celery cells and sink tissues
     ***protein***
   of celery, celeriac, and parsley subjected to sodium dodecyl sulfate gel
   electrophoresis showed a single major immuno-reactive 40-kD
     ***protein***
   ANSWER 11 OF 37 CAPLUS COPYRIGHT 2003 ACS
                       1996:111382 CAPLUS
 ESSION NUMBER:
                       124:141355
'UMENT NUMBER:
                       In vitro biosynthesis of caffeine; the stability of
                       N-methyltransferase activity in cell-free preparations
                       from liquid endosperm of Coffea arabica
                       Gillies, F. M.; Jenkins, G. I.; Ashihara, H.; Crozier,
'HOR(S):
                       Institute of Biomedical and Life Sciences, University
₹PORATE SOURCE:
                       of Glasgow, Glasgow, G12 8QQ, UK
                       Colloque Scientifique International sur le Cafe,
JRCE:
                        [Comptes Rendus] (1995), 16th(Vol. 2, Seizieme
                       Colloque Scientifique International sur le Cafe, 1995,
                       Vol. 2), 599-605
                       CODEN: CICRD8
                       Association Scientifique Internationale du Cafe
3LISHER:
CUMENT TYPE:
                       Journal
                       English
VGUAGE:
   A cell-free system has been developed from C. arabica that is a rich
   source of the N-methyltransferase activity which catalyzes the transfer of
   the Me group from S-adenosyl-L-methionine to methylxanthines producing
                ***Purifn*** . of the enzyme by ***anion***
                      ***chromatog*** . results in low activity yields.
     ***exchange***
   Part of the reason was found to be inhibition of the N-methyltransferase
   activity by KCl and NaCl which are used to ***elute***
     ***protein*** during ***anion*** - ***exchange***
     ***chromatog*** . The enzyme was found to have a half life at 4.degree.C
   of approx. 90 min. An extensive study of a wide range of protease
   inhibitors failed to show any effective stabilization of activity
   suggesting that the losses are not due to the action of endogenous
   proteases in the ext. The stability of the enzyme has been shown to be
   improved substantially with the incorporation of 20% (vol./vol.) glycerol
   or 20% (vol./vol.) ethylene ***glycol*** in the buffers used.
   Incorporation of 20% (vol./vol.) glycerol in buffers during ***anion***
                       ***chromatog*** . resulted in 54 - 78% yield of

    ***exchange***

   N-methyltransferase activity and a ca. 10-20 fold
                                                        ***purifn***
                                                        DUPLICATE 7
2 ANSWER 12 OF 37
                       MEDLINE
CESSION NUMBER:
                  95102513
                               MEDLINE
                             PubMed ID: 7804133
CUMENT NUMBER:
                  95102513
                  Rapid purification of tRNA(Lys) from rat liver.
TLE:
                  Kumar A M; Vulimiri S V; Nayak R
THOR:
                  Department of Microbiology and Cell Biology, Indian
RPORATE SOURCE:
                  Institute of Science, Bangalore.
                  BIOCHEMISTRY AND MOLECULAR BIOLOGY INTERNATIONAL, (1994
URCE:
                  Aug) 33 (6) 1081-9.
                  Journal code: 9306673. ISSN: 1039-9712.
JB. COUNTRY:
                  Australia
                  Journal; Article; (JOURNAL ARTICLE)
CUMENT TYPE:
MGUAGE:
                  English
                  Priority Journals
LE SEGMENT:
JTRY MONTH:
                  199501
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Entered STN: 19950215

Last Updated on STN: 19980206

ITRY DATE:

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Fast ***protein*** liquid pmatography (FPLC) system usin ono Q (HR 5/5) ***anion*** - ***e hange*** ***column***
  chromatography followed by highly cross-linked ***urea***
  -polyacrylamide gel electrophoresis ( ***urea*** -PAGE) was used for the
                         of lysine-specific tRNA (tRNA(Lys)) from rat liver.
    ***purification***
  Crude tRNA from rat liver was fractionated with a linear gradient of NaCl
  (0.3-0.8 M) in triethanolamine-HCl buffer, pH 4.5, and the activity of
  tRNA(Lys) was found to ***elute*** between 0.51 and 0.57 M NaCl. Using
  this concentration range of NaCl, tRNA(Lys) was refractionated on the same
  column with a shallow gradient, where a single peak of tRNA(Lys) activity
  was obtained. tRNA(Lys)-rich fractions recovered from the second run were
  electrophoretically separated on 16% polyacrylamide-7 M ***urea***
  into one major band and three minor bands. The major band showed a
  specific activity of 997 pmols/A260 U for tRNALys with a 43-fold
                        and approximately 17% recovery. The minor bands
     ***purification***
  displayed negligible or no activity for lysine. tRNA(Lys) obtained by this
  method was found to be homogeneous by competitive aminoacylation. The
  advantages of FPLC followed by ***urea*** -PAGE in the
    ***purification*** of an amino acid-specific tRNA over conventional
  column chromatography are discussed.
                                                       DUPLICATE 8
                      MEDLINE
2 ANSWER 13 OF 37
                               MEDLINE
                  95071402
CESSION NUMBER:
                            PubMed ID: 7980550
                  95071402
CUMENT NUMBER:
                  Rapid isolation of G alpha 13 from bovine brain membranes:
TLE:
                  supportive effect of ethylene glycol.
                  Harhammer R; Nurnberg B; Spicher K; Schultz G
THOR:
                  Institut fur Pharmakologie, Universitatsklinikum Rudolf
RPORATE SOURCE:
                  Virchow, Freie Universitat Berlin, Germany.
                  BIOCHEMICAL AND BIOPHYSICAL RESEARCH COMMUNICATIONS, (1994
URCE:
                  Oct 28) 204 (2) 835-40.
                  Journal code: 0372516. ISSN: 0006-291X.
                  United States
B. COUNTRY:
                  Journal; Article; (JOURNAL ARTICLE)
CUMENT TYPE:
                  English
NGUAGE:
                  Priority Journals
LE SEGMENT:
TRY MONTH:
                  199411
                  Entered STN: 19950110
TRY DATE:
                  Last Updated on STN: 20000303
                  Entered Medline: 19941130
   G13 belongs to the G12-subfamily of heterotrimeric regulatory G-
                                                           ***isolated***
     ***proteins*** . Employing specific antibodies, we
   alpha 13 from bovine brain by a four-step ***purification***
   combining conventional and affinity chromatography. The use of ethylene
                                                           ***elution***
     ***glycol*** as a protective agent influenced the
   properties of G alpha 13 markedly. Only in the presence of ethylene
     ***glycol*** (30% v/v) a clear separation of G alpha 13 from other G-
     ***proteins*** was achieved during the initial ***an
***exchange*** ***chromatography*** . This allowed
                                                        ***anion***
                                                               ***isolation***
   of G alpha 13 by subunit exchange chromatography on beta gamma-agarose. G
   alpha 13 was only released from immobilized beta gamma-dimers via
   activation by AMF but not by GTP gamma S, pointing to a poor basal
   nucleotide exchange of this ***protein*** . In contrast, N-terminally
   truncated G alpha 13 did not bind to immobilized beta gamma-dimers.
2 ANSWER 14 OF 37 CAPLUS COPYRIGHT 2003 ACS
                                                      DUPLICATE 9
                       1994:187267 CAPLUS
CESSION NUMBER:
                       120:187267
CUMENT NUMBER:
                       Sucrose-phosphate synthase is regulated via
TLE:
                       metabolites and protein phosphorylation in potato
                       tubers, in a manner analogous to the enzyme in leaves
                       Reimholz, Ralph; Geigenberger, Peter; Stitt, Mark
JTHOR(S):
                       Univ. Bayreuth, Bayreuth, D-95447, Germany
RPORATE SOURCE:
                       Planta (1994), 192(4), 480-8
CODEN: PLANAB; ISSN: 0032-0935
OURCE:
CUMENT TYPE:
                       Journal
                       English
NGUAGE:
                                          ***purified***
   Sucrose-phosphate synthase (SPS) was
                                                            40-fold from
   stored potato (Solanum tuberosum L.) tubers to a final specific activity
   of 33-70 nkat (mg ***protein*** )-1 via batch ***elution***
   diethylaminoethyl (DEAE) -sephacel, polyethylene ***glycol*** (PEG)
```

chromatog

Immunoblotting revealed a majo and a minor band with mol. wts 124.8 kba and 133.5 kDa, resp. Both ands were also present in exts. Frepd. in boiling SDS to exclude proteolysis. No smaller ***polypeptides*** were seen, except when the prepns. were incubated before application on a polyacrylamide gel. The enzyme prepn. was activated by glucose-6-phosphate and inhibited by inorg. phosphate. Both effectors had a large effect on the Km (fructose-6-phosphate) and the Km (uridine-5-diphosphoglucose) with phosphate acting antagonistically to glucose-6-phosphate. Preincubation of potato slices with low concns. of okadaic acid or microcystin resulted in a 3-4-fold decrease in the activity of SPS when the tissue was subsequently extd. and assayed. decrease was esp. marked when the assay contained low concns. of substrates and glucose-6-phosphate, and inorg. phosphate was included. Preincubation with mannose or in high osmoticum resulted in an increase of SPS activity. Analogous changes were obsd. in germinating Ricinus communis L. seedlings. After preincubation of the cotyledons in glucose, high SPS activity could be measured, whereas okadaic acid, omission of glucose, or addn. of phosphate or sucrose led to a large decrease of SPS activity in the selective assay. It is argued that SPS from non-photosynthetic tissues is regulated by metabolites and by ***protein*** phosphorylation in an analogous manner to the leaf enzyme.

DUPLICATE 10 MEDLINE 2 ANSWER 15 OF 37

MEDLINE PESSION NUMBER: 94163329

PubMed ID: 8118550 CUMENT NUMBER: 94163329

Use of polyethylene glycol and high-performance liquid CLE:

chromatography for preparative separation of Aspergillus

ficuum acid phosphatases.

Hamada J S THOR:

US Department of Agriculture, Southern Regional Research RPORATE SOURCE:

Center, New Orleans, LA 70179.

JOURNAL OF CHROMATOGRAPHY. A, (1994 Jan 14) 658 (2) 371-80. JRCE:

Journal code: 9318488.

Netherlands B. COUNTRY:

Journal; Article; (JOURNAL ARTICLE) CUMENT TYPE:

NGUAGE: English

LE SEGMENT: Priority Journals

TRY MONTH: 199404

TRY DATE: Entered STN: 19940412

> Last Updated on STN: 19940412 Entered Medline: 19940407

Proteins of Aspergillus ficuum culture filtrate were sequentially fractionated with 4, 9, 15, 19, 24, 30 and 36% polyethylene (PEG) into seven acid phosphatases (APases) with 93% and ***glycol*** 52% overall recoveries of activity and ***protein*** , respectively. Crude extract was also separated into seven APase peaks on a 30 cm x 2.5 ***column*** using 0.1 M ***anion*** - ***exchange*** Tris-HCl (pH 8.0) and a 0-0.4 M KCl gradient as the eluent, but their resolution was incomplete. However, when individual PEG precipitates were injected on to the column, each APase was ***eluted*** in a single, ***purification*** large peak resulting in 85% recovery and fifteen-fold of APase activity over the PEG precipitates. Use of PEG prior to HPLC separations also reduced the separation time to half and allowed a tenfold increase in sample load with complete resolution. The APases in PEG fractions and their corresponding HPLC peaks varied significantly in their kinetic parameters, including substrate specificity and pH optimum. The method developed is most beneficial for the ***isolation*** closely related APases from microbial or other sources for further molecular biology studies.

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.2 ANSWER 16 OF 37 CAPLUS COPYRIGHT 2003 ACS
                                                   DUPLICATE 11
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1994:161885 CAPLUS

120:161885

Semipreparative isolation of bovine casein components

by high-performance liquid chromatography

Ng-Kwai-Hang, K. F.; Chin, Dong

Dep. Anim. Sci., McGill Univ., Ste Anne de Bellevue,

QC, H9X 3V9, Can.

International Dairy Journal (1994), Volume Date 1993,

4(2), 99-110

CODEN: IDAJE6; ISSN: 0958-6946

CESSION NUMBER:

CUMENT NUMBER: [TLE:

JTHOR(S):

DRPORATE SOURCE:

OURCE:

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English
IGUAGE:
                                  ROTEIN*** PAK DEAE-15 HR
  An HPLC technique using a
                                    was developed for the
                                                              ***isolation***
                       ***column***
    ***exchange***
  of electrophoretically pure .kappa.-casein, .beta.-casein,
  .alpha.s2-casein, and .alpha.s1-casein from 250 mg of whole casein.
  mM tris buffer in 4.5 M ***urea*** at pH 7.0 and a convex NaCl
  gradient from 0 to 0.15 M, followed by a concave NaCl gradient from 0.15
  to 0.4 M provided the optimum conditions for sepn. of the casein
  components. The actual sepn. time of the 4 major caseins was less than 1
     After allowing time for column regeneration and equilibration, whole
  casein samples for sepn. could be loaded into the system every 2 h.
  Quant. measurements of the ***eluted*** fractions correlated well with
  known literature values and all the casein components were accounted for.
                                                     DUPLICATE 12
 ANSWER 17 OF 37
                      MEDLINE
                              MEDLINE
                94032311
CESSION NUMBER:
                           PubMed ID: 8218256
                 94032311
CUMENT NUMBER:
                 Secondary structure analysis of purified functional CHIP28
TLE:
                 water channels by CD and FTIR spectroscopy.
                 Van Hoek A N; Wiener M; Bicknese S; Miercke L; Biwersi J;
THOR:
                 Verkman A S
                 Department of Medicine, University of California, San
RPORATE SOURCE:
                 Francisco 94143.
NTRACT NUMBER:
                 DK35124 (NIDDK)
  DK43840 (NIDDK)
  HL42368 (NHLBI)
                 BIOCHEMISTRY, (1993 Nov 9) 32 (44) 11847-56.
URCE:
                 Journal code: 0370623. ISSN: 0006-2960.
                 United States
B. COUNTRY:
                 Journal; Article; (JOURNAL ARTICLE)
CUMENT TYPE:
NGUAGE:
                 English
                 Priority Journals
LE SEGMENT:
                 199312
TRY MONTH:
                 Entered STN: 19940117
TRY DATE:
                 Last Updated on STN: 19970203
                 Entered Medline: 19931216
                                          CHIP28 is an important water
  The integral membrane ***protein***
  channel in erythrocytes and kidney tubule epithelia and is a member of a
  family of channel/pore ***proteins*** including the lens
                   MIP26. The purposes of this study were to
    ***protein***
  functional, delipidated CHIP28 to homogeneity and to determine secondary
  structure by circular dichroism (CD) and Fourier transform infrared
  spectroscopy (FTIR). CHIP28 was initially ***purified***
                  ***anion*** - ***exchange***
                                                      ***chromatography***
  delipidated by
  following solubilization of N-lauroylsarcosine-stripped erythrocyte
  membranes with beta-octylglucoside (OG); MIP26 was initially
                                         ***anion*** - ***exchange***
                    and delipidated by
     ***purified***
                          following solubilization of
                                                        ***urea*** -stripped
     ***chromatography***
  bovine lens membranes by monomyristoylphosphatidylcholine. CHIP28
   (glycosylated and nonglycosylated) and MIP26 were ***purified***
   further by high-performance size-exclusion chromatography,
   in OG as apparent dimers and tetramers, respectively. Proteoliposomes
                      ***purified*** CHIP28 were highly water-permeable,
   reconstituted with
  with an osmotic water permeability Pf of 0.04 cm/s at 10 degrees C that
  was inhibited by 0.1 mM HgCl2. Proteoliposomes reconstituted with MIP26
  had a low Pf of 0.005 cm/s. CD spectra of CHIP28 in OG or in reconstituted
  proteoliposomes gave a maximum at 193 nm and minima at 208 and 222 nm.
   Spectral decomposition using ***protein*** basis spectra gave 40 +/-
   5% alpha-helix and 43 +/- 3% beta-sheet and -turn. HgCl2 did not affect
   the CD spectrum of CHIP28. Attenuated total reflectance FTIR of air-dried,
   membrane-associated CHIP28 gave 38 +/- 5% alpha-helix and 40 +/- 4%
   beta-sheet and -turn by spectral decomposition of the amide I resonance.
   For comparison, CD of MIP26 in OG gave 49 +/- 7% alpha-helix and 32 +/-
   12% beta-sheet and -turn; FTIR gave 32 +/- 8% alpha-helix and 45 +/- 6%
   beta-sheet and -turn. Analysis of CHIP28 and MIP26 sequence data by the
   generalized hydropathy method of Jahnig [Jahnig, F. (1990) Trends Biochem.
   Sci. 15, 93-95] predicted 39-47% alpha-helix and 15-20% beta-structures.
   These results establish procedures to obtain large quantities of pure
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CHIP28 and MIP26 in functional forms and provide evidence for multiple

membrane-spanning alpha-helices or mixed alpha/beta-domains.

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ANSWER 18 OF 37 CAPLUS COPYR T 2003 ACS
                                                  DUPLICATE 13
CESSION NUMBER: 1993:54851
                                  APLUS
                     118:54851
CUMENT NUMBER:
                     Purification and characterization of
CLE:
                     1-aminocyclopropane-1-carboxylate N-malonyltransferase
                     from etiolated mung bean hypocotyls
                     Guo, Lining; Arteca, Richard N.; Phillips, Allen T.;
THOR(S):
                     Liu, Yu
                     Dep. Hortic., Pennsylvania State Univ., University
RPORATE SOURCE:
                     Park, PA, 16802, USA
                      Plant Physiology (1992), 100(4), 2041-5
JRCE:
                      CODEN: PLPHAY; ISSN: 0032-0889
                      Journal
CUMENT TYPE:
VGUAGE:
                      English
  1-Aminocyclopropane-1-carboxylate (ACC) N-malonyltransferase converts ACC,
  an immediate precursor of ethylene, to the presumably inactive product
  malonyl-ACC (MACC). This enzyme plays a role in ethylene prodn. by
  reducing the level of free ACC in plant tissue. In this study, ACC
                                             3660-fold from etiolated mung
  N-malonyltransferase was ***purified***
  bean (Vigna radiata) hypocotyls, with a 6% overall recovery. The final
  specific activity was about 83,000 nmol of MACC formed mg-1
  ***protein*** h-1. The five-step ***purifn*** . protocol consisted of polyethylene ***glycol*** fractionation, Cibacron blue 3GA-agarose
  chromatog. using salt gradient ***elution*** , Sephadex G-100 gel
                     ***anion*** - ***exchange***
                                                        ***chromatog***
  filtration, MonoQ
  and Cibacron blue 3GA-agarose chromatog. using malonyl-CoA plus ACC for
    ***elution*** . The mol. mass of the native enzyme detd. by Sephadex
  G-100 chromatog. was 50 kD. ***Protein*** from the final
    ***purifn***^{-} . step showed one major band at 55 kD after sodium dodecyl
  sulfate polyacrylamide gel electrophoresis, indicating that ACC
  N-malonyltransferase is a monomer. The mung bean ACC N-malonyltransferase
  has a pH optimum of 8.0, an apparent Km of 0.5 mM for ACC and 0.2 mM for
  malonyl-CoA, and an Arrhenius activation energy of 70.29 kJ mol-1
  degree-1.
                                                     DUPLICATE 14
                      MEDLINE
2 ANSWER 19 OF 37
CESSION NUMBER: 92201243
                             MEDLINE
                           PubMed ID: 1839382
                 92201243
CUMENT NUMBER:
                 Identification of the microvillar 110-kDa calmodulin
TLE:
                 complex (myosin-1) in kidney.
                 Coluccio L M
THOR:
                 Department of Biochemistry, Emory University School of
RPORATE SOURCE:
                 Medicine, Atlanta, GA 30322.
                 1RO1 GM44211 (NIGMS)
NTRACT NUMBER:
  S07 RR05364 (NCRR)
                 EUROPEAN JOURNAL OF CELL BIOLOGY, (1991 Dec) 56 (2) 286-94.
URCE:
                 Journal code: 7906240. ISSN: 0171-9335.
                 GERMANY: Germany, Federal Republic of
B. COUNTRY:
                 Journal; Article; (JOURNAL ARTICLE)
CUMENT TYPE:
                 English
NGUAGE:
                 Priority Journals
LE SEGMENT:
                 199204
ITRY MONTH:
                 Entered STN: 19920509
ITRY DATE:
                 Last Updated on STN: 19920509
                 Entered Medline: 19920430
   The epithelial layer lining the proximal convoluted tubule of mammalian
   kidney contains a brush border of numerous microvilli. These microvilli
   appear in structure to be very similar to the microvilli on epithelial
   cells of the small intestine. Microvilli found in both the small intestine
   and the proximal convoluted tubules in kidney have a core bundle of actin
   filaments bundled by the accessory ***proteins*** villin and fimbrin.
   Along the length of intestinal microvilli, lateral links can be observed
   to connect the core bundle of actin filaments to the membrane. These
   cross-bridges are comprised of a 110-kDa calmodulin complex which belongs
   to a class of single-headed myosin molecules, collectively referred to as
   myosin-1. We now report that an analogous calmodulin-binding
     ***polypeptide*** of 105 kDa has been identified in rat kidney cortex.
                                   is preferentially found in
                ***polypeptide***
     ***purified*** kidney brush borders, can be extracted with ATP, and co-
     ***elutes*** with calmodulin on gel filtration and ***anion***
```

madeat Ambage activity in buffer

containing CaCl2. The partiall ***purified*** 105-kDa ***polypeptide*** will bin lodinated calmodulin and will s Ament with F-actin in buffer containing ethylene ***glycol*** -bis-(betaaminoethyl ether)-N,N,N',N'-tetraacetic acid (EGTA) or Ca2+. The addition of ATP partially reverses this association with F-actin. These results indicate that myosin-1, in addition to its presence in intestinal brush borders, is present in the brush border of kidney. We also provide ***polypeptide*** preliminary evidence to indicate that the 105-kDa not restricted to tissues possessing a brush border. **DUPLICATE 15** 2 ANSWER 20 OF 37 MEDLINE CESSION NUMBER: 92338871 MEDLINE CUMENT NUMBER: 92338871 PubMed ID: 1668267 High-yield purification of potato tuber pyrophosphate: TLE: fructose-6-phosphate 1-phosphotransferase. Moorhead G B; Plaxton W C THOR: Department of Biology, Queen's University, Kingston RPORATE SOURCE: Ontario, Canada. PROTEIN EXPRESSION AND PURIFICATION, (1991 Feb) 2 (1) URCE: 29-33. Journal code: 9101496. ISSN: 1046-5928. United States JB. COUNTRY: Journal; Article; (JOURNAL ARTICLE) CUMENT TYPE: MGUAGE: English LE SEGMENT: Priority Journals ITRY MONTH: 199208 Entered STN: 19920911 ITRY DATE: Last Updated on STN: 19920911 Entered Medline: 19920826 The procedure of Yuan et al. (1988, Biochem. Biophys. Res. Commun. 154, ***isolation*** of potato pyrophosphate:fructose-6-

111-117) for the phosphate 1-phosphotransferase (PFP) has been modified so that a high yield of homogeneous enzyme could be obtained. Modifications included a lower temperature heat step, a lower percentage initial polyethylene fractionation step (0 to 4%, w/v), stepwise ***elution*** ***glycol***

following an increase from 30 to 50 mM pyrophosphate during affinity ***anion*** chromatography on Whatman P11 phosphocellulose,

using Q-Sepharose "Fast Flow," ***chromatography*** ***exchange*** and gel filtration chromatography with Superose 6 "Prep grade." Our procedure resulted in an overall 42% yield and a final specific activity of 87 mumol fructose 1,6-bisphosphate produced per minute per milligram

protein . Rabbit anti-(potato PFP) polyclonal antibodies effectively immunoprecipitated the activity of both the pure enzyme and the enzyme from a crude extract. Western blot analysis demonstrated that the antibodies were monospecific for PFP. A survey of various potato cultivars demonstrated significant differences in PFP activity with respect to fresh weight. This observation should be taken into ***purification*** of potato PFP is consideration before any undertaken.

DUPLICATE 16 .2 ANSWER 21 OF 37 MEDLINE

CESSION NUMBER: 91009234 MEDLINE

CUMENT NUMBER: 91009234 PubMed ID: 2170394

Purification and characterization of glycosyl-TLE: phosphatidylinositol-specific phospholipase D.

Huang K S; Li S; Fung W J; Hulmes J D; Reik L; Pan Y C; Low JTHOR:

RPORATE SOURCE: Department of Protein Biochemistry, Hoffmann-La Roche Inc.,

Nutley, New Jersey 07110.

GM-35873 (NIGMS) NTRACT NUMBER:

GM-40083 (NIGMS)

CUMENT TYPE:

JOURNAL OF BIOLOGICAL CHEMISTRY, (1990 Oct 15) 265 (29) URCE:

17738-45.

Journal code: 2985121R. ISSN: 0021-9258.

JB. COUNTRY: United States

Journal; Article; (JOURNAL ARTICLE)

MGUAGE: English

Priority Journals LE SEGMENT:

TRY MONTH: 199011

Entered STN: 19910117 TRY DATE:

Last Updated on STN: 19970203 Entered Medline: 19901121

We have developed a simple immunaffinity chromatography proced for the state of a glyssyl-phosphatidylinositol (GPI) pecific phospholipase D (GPI-PLD) from bovine serum. The enzyme was initially ***purified*** by a procedure consisting of 9% polyethylene ***glycol*** precipitation, Q Sepharose ***anion*** - ***exchange*** ***chromatography*** , S-300 gel filtration, wheat germ lectin-Sepharose, hydroxylapatite agarose, zinc chelate matrix, Mono Q-high performance liquid chromatography (HPLC), and Superose 12 (gel filtration) HPLC. Using ***purified*** material as immunogen, we generated a panel of monoclonal antibodies. A low affinity antibody was selected for the ***purification*** of catalytically active GPI-PLD from bovine serum by immunoaffinity chromatography, followed by wheat germ lectin-Sepharose and Mono Q-fast ***protein*** liquid chromatography. The latter method provides a simple ***purification*** procedure with an overall yield of 26%. The ***purified*** enzyme has an apparent molecular weight of about 100,000 by sodium dodecyl sulfate-polyacrylamide gel electrophoresis and a pI of about 5.6 by isoelectric focusing gel analysis. On Superose 12 HPLC, the material ***purified*** by the latter method ***elutes*** as a single peak with an apparent molecular weight of 200,000 as determined by ***protein*** standards. The enzyme activity is inhibited by [ethylenebis(oxyethylenenitrilo)]tetraacetic acid or 1,10-phenanthroline. Phosphatidic acid is the only 3H-labeled product when [3H] myristate-labeled variant surface glycoprotein is hydrolyzed by the ***purified*** enzyme. Amino terminal sequence analysis of the intact 100-kDa ***protein*** reveals no strong homology to that of any other known ***protein***. Twelve tryptic ***peptides*** derived from the intact ***protein*** have been subjected to amino acid sequence analysis. Two of them share sequence homology with each other and with the metal ion binding domains of members of the integrin family. Based upon these criteria, it appears that the ***purified*** enzyme is distinct from other phospholipases with specificity for inositol phospholipids.

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DUPLICATE 17
 ANSWER 22 OF 37 CAPLUS COPYRIGHT 2003 ACS
                      1991:160378 CAPLUS
CESSION NUMBER:
                       114:160378
CUMENT NUMBER:
                       Two class I aldolases in Klebsormidium flaccidum
                        (Charophyceae): an evolutionary link from
                       chlorophytes to higher plants
                       Jacobshagen, Sigrid; Schnarrenberger, Claus
THOR (S):
                       Inst. Pflanzenphysiol. Mikrobiol., Freie Univ. Berlin,
RPORATE SOURCE:
                       Berlin, D-1000/33, Germany
                        Journal of Phycology (1990), 26(2), 312-17
JRCE:
                        CODEN: JPYLAJ; ISSN: 0022-3646
CUMENT TYPE:
                        Journal
NGUAGE:
                       English
  Two fructose-diphosphate aldolase from K. flaccidum were
                                                                ***purified***
   by affinity ***elution*** from phosphocellulose. The 2 enzymes were
   subsequently sepd. by HPLC on an ***anion*** - ***exchange***
***column*** (QAE-silica). The aldolase ***eluting*** fi
   represented 5% of the total activity; the other aldolase represented the
   remaining activity. The activity of the enzymes was not reduced by the
   presence of 1 mM EDTA or increased by 0.1 mM Zn2+, establishing their
  character as class I type (Me2+ independent) aldolases. The Km values were 1.7 and 34.7 .mu.M for the enzyme ***eluting*** first and second,
   resp., from the QAE-silica column. The subunit mol. masses, as detd. by
   SDS-PAGE, were 40.5 and 37 kD; the specific activities of the
     ***purified*** enzymes were 7.9 and 24.7 U/mg ***protein***
   The 2 aldolases of K. flaccidum are homologous to the cytosol and
   chloroplast specific isoenzymes of higher plants by several criteria and
   are therefore probably located in the same cellular compartments in K.
   flaccidum. The Km and specific activity for the chloroplast aldolase of
   K. flaccidum are 3 times higher than for the chloroplast aldolase of
   higher plants, a remarkable difference. Immunotitrn. with specific
   antisera against the chloroplast aldolase of Chlamydomonas reindardtii and
   spinach showed that the chloroplast aldolase of K. flaccidum was
   immunochem. intermediate in structure to the resp. aldolases of C.
   reinhardtii and higher plants. K. flaccidum is the second species of
   Charophyceae (besides Chara foetida) with 2 class I aldolases as in higher
   plants whereas 2 species of Chorophyceae have only one class I aldolase
   and, under some conditions, an addnl. class II (Me2+ dependent) aldolase.
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Thus, aldolases may turn out, in addn. to the known enzymes of glycolate

to be a noticel engume

system to evaluate algal evolution along with cytol. features.

ANSWER 23 OF 37 CAPLUS COPYRIGHT 2003 ACS SSION NUMBER: 1990:104838 CAPLUS 112:104838 MENT NUMBER: Process for the chromatographic purification of a E: 69,000-dalton outer membrane protein of Bordetella pertussis for vaccines Burns, D. L.; Brennan, M. J.; Gould-Kostka, J. L.; NTOR(S): Manclark, C. R. United States Dept. of Health and Human Services, USA :NT ASSIGNEE(S): U. S. Pat. Appl., 14 pp. Avail. NTIS Order No. ≀CE: PAT-APPL-7-308 864. CODEN: XAXXAV MENT TYPE: Patent English BUAGE:

ILY ACC. NUM. COUNT: 2

ENT INFORMATION:

PATENT NO. KIND DATE APPLICATION NO. DATE US 308864 A0 19890715 US 5101014 A 19920331 US 1989-308864 19890210 US 1989-308864 19890210 ITY APPLN. INFO.: US 1989-308864 19890210
The title process involves prepg. a ***protein*** ext. contg. the ORITY APPLN. INFO.: ***protein*** , applying the ext. to an ***anion*** - ***exchange***
column (e.g. DEAE-Sepharose), sepg. the ***protein*** from th ***eluting*** it with a linear salt gradient, pooling fractions contg. the ***protein*** , applying the pooled fractions to an affinity column contg. a ***protein*** -specific binding medium (e.g. Affi-Gel Blue), and ***eluting*** the ***purified*** ***protein*** (e.g. with .apprx.4 M ***urea***). The
protein was ***purified*** from a ***protein*** ext. of B. pertussis Bp 353. It bound monoclonal antibody BPE3.

DUPLICATE 18 2 ANSWER 24 OF 37 MEDLINE CESSION NUMBER: 89109147 MEDLINE CUMENT NUMBER: 89109147 PubMed ID: 2521480 Analysis of the proteoglycans synthesized by corneal CLE:

explants from embryonic chicken. II. Structural characterization of the keratan sulfate and dermatan

sulfate proteoglycans from corneal stroma.

Midura R J; Hascall V C THOR:

RPORATE SOURCE: Proteoglycan Chemistry Section, National Institute of

Dental Research, Bethesda, Maryland 20892.

NTRACT NUMBER: EY05779 (NEI)

JOURNAL OF BIOLOGICAL CHEMISTRY, (1989 Jan 25) 264 (3) URCE:

1423-30.

Journal code: 2985121R. ISSN: 0021-9258.

B. COUNTRY: United States

CUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

NGUAGE: English

LE SEGMENT: Priority Journals

ITRY MONTH: 198903

ITRY DATE: Entered STN: 19900308

> Last Updated on STN: 19970203 Entered Medline: 19890303

Radioisotopically labeled proteoglycans were ***isolated*** from a 4 M ***guanidine*** HCl, 2% Triton X-100 extract of corneal stroma from day

18 chicken embryos by ***anion*** - ***exchange***

chromatography . Two predominant proteoglycans in the sample were separated by octyl-Sepharose chromatography using a gradient

elution of detergent in 4 M ***guanidine*** proteoglycan had an overall mass of approximately 125 kDa, a single dermatan sulfate chain (approximately 85-90% chondroitin 4-sulfate, low iduronate content) of approximately 65 kDa, and a core ***protein*** after chondroitinase ABC digestion of approximately 45 kDa which also contained one to three N-linked oligosaccharides and one O-linked oligosaccharide. The other proteoglycan had an overall size of approximately 100 kDa, two to three keratan sulfate chains of approximately 100 kDa, two to thick here in the following approximately 15 kDa each, and a core ***protein*** following

N-linked but no O-linked oligo charides. A larger size, a grecoverall hydrophobicity (as measured by its interaction with octyl-Sepharose) and an absence of O-linked oligosaccharides argue that this core ***protein*** is a distinct gene product from the core of the dermatan sulfate proteoglycan. ***protein*** **DUPLICATE 19** 2 ANSWER 25 OF 37 MEDLINE MEDLINE CESSION NUMBER: 88100442 PubMed ID: 3501173 88100442 CUMENT NUMBER: Partial purification and characterization of extrinsic TLE: pathway inhibitor (the factor Xa-dependent plasma inhibitor of factor VIIa/tissue factor). Warn-Cramer B J; Maki S L; Zivelin A; Rapaport S I THOR: Department of Medicine, University of California, San Diego RPORATE SOURCE: Medical Center 92103. HL 07107 (NHLBI) NTRACT NUMBER: HL 27234 (NHLBI) THROMBOSIS RESEARCH, (1987 Oct 1) 48 (1) 11-22. URCE: Journal code: 0326377. ISSN: 0049-3848. United States B. COUNTRY: Journal; Article; (JOURNAL ARTICLE) CUMENT TYPE: English NGUAGE: Priority Journals LE SEGMENT: 198802 TRY MONTH: Entered STN: 19900305 TRY DATE: Last Updated on STN: 20000303 Entered Medline: 19880202 partially from plasma We report a procedure to ***purify*** (approximately 1200 fold) the factor Xa-dependent inhibitor of ***VIIa*** /tissue factor (i.e., the extrinsic pathway ***factor*** inhibitor or EPI) and describe some of its properties. An assay for EPI was developed based upon inhibition of ***factor*** ***VIIa*** /tissue factor induced release of activation ***peptide*** tritiated factor IX by a test sample in the presence but not in the absence of factor Xa. Approximately 50% of the total EPI activity in plasma was found in the lipoprotein fraction, which was used as the starting material for ***purification*** . Total lipoproteins (***urea*** soluble apoproteins gel filtered on Sephacryl S-200. The inhibitory activity co- ***eluted*** with the major ***protein*** peak, which primarily contained apoprotein A-I. Inhibitory activity was separated from apoprotein A-I by ***anion*** - ***exchange***

isolated by density ultracentrifugation) were delipidated and the ***chromatography*** on Q-Sepharose and was further resolved from higher

and lower molecular weight contaminating ***proteins*** by polypreparative disc gel electrophoresis in the presence of 0.1% SDS. Functional inhibitory activity ***eluted*** from the polypreparative disc gel in two discrete pools of different molecular weights (approximately 34,000 and approximately 43,000 D). Apoprotein E was identified by immunological techniques as the major ***protein*** present in both of these pools. However, incubation with a monospecific polyclonal antibody to human apoprotein E did not decrease EPI activity either in plasma or in the partially ***purified*** polypreparative disc gel fractions. A rabbit antiserum was prepared against material from the polypreparative disc gel. The IgG fraction neutralized approximately 95% of the total inhibitory activity present in plasma. Therefore, EPI in the lipoprotein fraction and in the non-lipoprotein fraction of plasma appears to be antigenically similar.

L2 ANSWER 26 OF 37 CAPLUS COPYRIGHT 2003 ACS 1993:401756 CAPLUS CESSION NUMBER: 119:1756 CUMENT NUMBER: Chromatographic purification of erythropoietin TLE: Por-Hsiung, Lai; Strickland, Thomas Wayne IVENTOR(S): Kirin-Amgen, Inc., USA ATENT ASSIGNEE(S): PCT Int. Appl., 20 pp. OURCE: CODEN: PIXXD2 CUMENT TYPE: Patent

English

AMILY ACC. NUM. COUNT: 1 ATENT INFORMATION:

ANGUAGE:

```
WO 1986-US1342 19860620
                 A1
                      1986123
     W: AU, DK, JP
     RW: AT, BE, CH, DE, FR, GB, IT, LU, NL, SE
                     19870519 US 1985-747119 19850620
  US 4667016
                A
                                                  19860420
                A1 19920621
                                  IL 1986-79176
  IL 79176
                                                  19860618
                A1 19920317
                                  CA 1986-511855
  CA 1297635
                                   ZA 1986-4573
                                                   19860619
                     19870225
  ZA 8604573
                                   ES 1986-556257
                                                   19860619
                 A1 19880101
  ES 556257
                                   EP 1986-904556
                                                   19860620
  EP 228452
                 A1 19870715
                 B1 19950322
     R: AT, BE, CH, DE, FR, GB, IT, LI, LU, NL, SE
                                  JP 1986-503570
                                                  19860620
  JP 63503352 T2 19881208
                B4 19941207
  JP 06098019
                                   AU 1986-61230
                                                  19860620
                B2 19910214
  AU 606578
                A1 19870113
  AU 8661230
                                   IL 1986-97135
                                                  19860620
  IL 97135
                A1 19920621
                                   AT 1986-904556 19860620
                E 19950415
  AT 120208
                                                  19870218
                A 19870218
                                  DK 1987-813
  DK 8700813
  CA 1312994
                                   CA 1991-616009 19910221
                 A2 19930119
                                 US 1985-747119
                                                   19850620
[ORITY APPLN. INFO.:
                                                 19860613
                                 US 1986-872152
                                 CA 1986-5118557
                                                  19860618
                                 IL 1986-79176
                                                   19860620
                                 WO 1986-US1342
                                                   19860620
        ***anion*** - ***exchange***
                                       ***chromatog*** . on
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Methods for chromatog. ***purifn*** . of erythropoietin from a variety of sources, including biol. fluids or transgenic animal cell lines, is described. The first method is a reversed-phase chromatog. that involves adsorption of the erythropoietin onto a C4 or C6 resin followed by ***elution*** with increasing concns. of EtOH (either stepwise or in a gradient); after removal of EtOH, an erythropoietin fraction of high specific activity with yield .gtoreq.50% is obtained. A second method DEAE-agarose at acid pH under conditions that prevent activation of acid proteinases is also described. The two methods may be combined for rapid ***purifn*** . of erythropoietin in high yield and purity. Culture supernatants from CHO cells stably expressing the erythropoietin gene on the plasmid pDSVL-gHuEPO were concd. by diafiltration and fractionated by chromatog. on VYDAC 214TP-B using a 0-80% EtOH gradient in 10 mM tris pH 7.0. The peak of UV absorption ***eluting*** around 60% EtOH was pooled and applied to a DEAE-agarose column which was washed with an acid ***urea*** buffer to remove proteinases and the ***urea*** removed and the column brought to neutral pH with a low-salt buffer. CuSO4 is optionally present in the wash to assist in oxidn. of sulfhydryl groups of undesired ***protein*** . Erythropoietin was ***eluted*** with a buffer contg. NaCl 75 mM. Final purity of the erythropoietin is >95% and is low in pyrogens and serum ***proteins***

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CESSION NUMBER:
                       1985:537917 CAPLUS
CUMENT NUMBER:
                       103:137917
                       Purification of a neurotrophic factor for ciliary
TLE:
                       neurons from chick intraocular tissue using
                       nondenaturing conditions
                       Watters, Diane; Hendry, Ian
JTHOR (S):
                        John Curtin Sch. Med. Res., Aust. Natl. Univ.,
PROPRATE SOURCE:
                        Canberra, Australia
                        Biochemistry International (1985), 11(2), 245-53
DURCE:
                        CODEN: BIINDF; ISSN: 0158-5231
CUMENT TYPE:
                        Journal
ANGUAGE:
                       English
```

2 ANSWER 27 OF 37 CAPLUS COPYRIGHT 2003 ACS

A neurotrophic factor for the parasympathetic ciliary ganglion neurons was ***purified*** to apparent homogeneity from chicken intraocular tissues by high-performance ***anion*** - ***exchange*** ***chromatog***. under nondenaturing conditions. The initial stages of ***purifn***. were carried out as described by G. Barbin et al. (1984) up to and including chromatog. on DEAE-cellulose. The 0.25M NaCl eluate from the DEAE-cellulose column was concd. by ultrafiltration, dild., reconcd., and subjected to fast- ***protein*** liq. chromatog. on a Pharmacia MonoQ column with gradient ***elution***. The active fractions were pooled, concd., radiolabeled with 125I, and subjected to electrophoresis on SDS-

```
of approx. 43,000 daltons and s not appear to be related to neurotrophic factor ***isola *** from the same source by
                                                                          parative
  SDS-gel electrophoresis. The final sp. activity of the ***isolated***
  neurotrophic factor was estd. to be .apprx.4 .times. 105 units/mg.
                                                          DUPLICATE 20
2 ANSWER 28 OF 37
                        MEDLINE
                 86026398
                                 MEDLINE
CESSION NUMBER:
                             PubMed ID: 3931690
                   86026398
CUMENT NUMBER:
                   Simultaneous purification of multiple forms of rat liver
                   microsomal cytochrome P-450 by high-performance liquid
                   chromatography.
                   Funae Y; Imaoka S
THOR:
                   BIOCHIMICA ET BIOPHYSICA ACTA, (1985 Oct 17) 842 (2-3)
URCE:
                   119-32.
                   Journal code: 0217513. ISSN: 0006-3002.
                   Netherlands
                   Journal; Article; (JOURNAL ARTICLE)
```

B. COUNTRY:

CUMENT TYPE:

NGUAGE: English

Priority Journals LE SEGMENT: TRY MONTH: 198511

Entered STN: 19900321 TRY DATE:

Last Updated on STN: 19970203

Entered Medline: 19851127

14 microsomal cytochromes P-450 were ***purified*** from the liver of untreated and phenobarbital- or 3-methylcholanthrene-treated male rats. Following solubilization of microsomes with sodium cholate, poly(ethylene ***glycol***) fractionation and aminohexyl-Sepharose 4B chromatography, cytochromes P-450 were ***purified*** by high-performance liquid chromatography (HPLC), using a preparative DEAE- ***anion*** ***column*** . The pass-through fraction was further ***exchange*** by HPLC using a cation-exchange column. Other fractions ***purified*** ***eluted*** on preparative DEAE-HPLC were further applied onto an HPLC using a DEAE-column. Five kinds (P-450UT-2-6), four kinds (P-450PB-1,2,4 and 5) and five kinds (P-450MC-1-5) of cytochromes P-450 were

purified from untreated rats or rats treated with phenobarbital or 3-methylcholanthrene, respectively. HPLC profiles of tryptic ***peptides*** of cytochromes P-450UT-2 and P-450MC-2 were identical and the other profiles obtained from seven ***purified*** cytochromes

P-450 were distinct from each other. Amino-terminal sequences of eight forms of cytochrome P-450 (UT-2, UT-5, PB-1, PB-2, PB-4, PB-5, MC-1 and MC-5) were distinct except for cytochromes P-450PB-4 and P-450PB-5.

2 ANSWER 29 OF 37 CAPLUS COPYRIGHT 2003 ACS 1984:171197 CAPLUS CESSION NUMBER:

100:171197 CUMENT NUMBER:

Partially purified osteogenic factor from TLE:

demineralized bone

Seyedin, Saeid; Thomas, Thomas VENTOR(S):

Collagen Corp., USA ATENT ASSIGNEE(S):

U.S., 7 pp. OURCE: CODEN: USXXAM Patent

CUMENT TYPE: English ANGUAGE:

AMILY ACC. NUM. COUNT: 1

ATENT INFORMATION:

MI INI ORIEITEON	•			
PATENT NO.		DATE	APPLICATION NO.	DATE
US 4434094	A	19840228	US 1983-484286 EP 1984-300322	
EP 121976 EP 121976	A3	19860723		10040110
CA 1223199	A1	19870623	CA 1984-446151	
JP 59190919 JP 63016364	A2	19841029	JP 1984-14983	19840130
OPTTV ADDIN TN	FO :		US 1983-484286 ***purified***	19830412 osteogenic factor
from deminera	lized mam	malian. e.g.	, cattle, pig, bone oteins*** were ex	particles is
demineralized	bone wit	h a dissocia	tive extractant, e. the presence of a p	g., 8M ***urea***
O1 11.			1 27 1 1 4	ide Who oved

N athulmaleimide The extd

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nonfibrous ***proteins*** were fractionated by ***anion**;

***exchange*** ***chroma *** . on DEAE-cellulose at pH
   fraction not adsorbed by the DEAE-cellulose was further fractionated by
   cation-exchange chromatog. on CM-cellulose at pH 4.8. The fraction
  adsorbed by CM-cellulose was ***eluted*** with a NaCl gradient, and the partially ***purified*** osteogenic factor was sepd. by gel
   chromatog. on Sephacryl S 200 as a ***protein*** with mol. wt.
   .ltoreq.30,000 daltons. The osteoinductive activity of this
                    was assayed by measuring its ability to stimulate rat
     ***protein***
   muscle fibroblasts in agarose to synthesize type II collagen and cartilage
   proteoglycans.
                                                       DUPLICATE 21
                       MEDLINE
2 ANSWER 30 OF 37
CESSION NUMBER: 84203512 MEDLINE
                            PubMed ID: 6232947
CUMENT NUMBER:
                 84203512
                 Adenosinetriphosphatase site stoichiometry in sarcoplasmic
TLE:
                  reticulum vesicles and purified enzyme.
                 Barrabin H; Scofano H M; Inesi G
THOR:
NTRACT NUMBER: HL 27867 (NHLBI)
                BIOCHEMISTRY, (1984 Mar 27) 23 (7) 1542-8.
URCE:
                  Journal code: 0370623. ISSN: 0006-2960.
B. COUNTRY:
                  United States
                  Journal; Article; (JOURNAL ARTICLE)
CUMENT TYPE:
MGUAGE:
                 English
                 Priority Journals
LE SEGMENT:
                  198406
ITRY MONTH:
                  Entered STN: 19900319
ITRY DATE:
                  Last Updated on STN: 19970203
                  Entered Medline: 19840622
   The stoichiometry of phosphorylation (catalytic) sites in sarcoplasmic
   reticulum vesicles ( SRV ) and SR ATPase ***purified***
   differential solubilization with deoxycholate was found to be 4.77 + /- 0.4
   and 6.05 + /- 0.18 \text{ nmol/mg of} ***protein*** , respectively, when
   phosphorylation was carried out under conditions permitting 32P labeling
   of nearly all sites. Assuming that each site corresponds to a single 115K
   ATPase chain, the observed site stoichiometry accounts only for 55% and
   70% of the total ***protein*** . Failure to obtain higher
   phosphorylation levels was due to the presence of nonspecific
     ***protein*** contaminants in SRV or to the presence of inactive
   aggregates in the ATPase ***purified*** with deoxycholate. This was
   demonstrated by dissolving SRV and ***purified*** ATPase with lithium
   dodecyl sulfate, subjecting them to molecular sieve HPLC, and collecting
         ***elution*** fractions for determination of ***protein***
   measurement of 32P-labeled sites, and electrophoretic analysis. In fact,
   in the specific ***elution*** peak containing the 115K ATPase chains,
   phosphorylation levels were 6.62 + /- 0.33 and 7.\overline{03} + /- 0.18 in SRV and
     ***purified*** ATPase, corresponding to 68% and 86% of the
     ***protein*** in the specific ***elution*** peak. An alternate
     ***purification*** method was then developed, based on solubilization of
   SRV with dodecyl octaethylene ***glycol*** monoether ( C12E8 ),
   separation of delipidated ATPase by ***anion*** - ***exchange***
     ***chromatography*** , and enzyme reactivation with phosphatidylcholine.
   This preparation yields 7.3 +/- 0.44 nmol of phosphorylation site/mg of
     ***protein*** of the SRV fraction before HPLC. (ABSTRACT TRUNCATED AT 250
   WORDS)
                                                       DUPLICATE 22
L2 ANSWER 31 OF 37
                      MEDLINE
CCESSION NUMBER: 84202839
                               MEDLINE
                            PubMed ID: 6721830
                  84202839
CUMENT NUMBER:
                  Studies on the alpha-subunit of bovine brain S-100 protein.
[TLE:
                  Masure H R; Head J F; Tice H M
BIOCHEMICAL JOURNAL, (1984 Mar 15) 218 (3) 691-6.
JTHOR:
OURCE:
                  Journal code: 2984726R. ISSN: 0264-6021.
JB. COUNTRY:
                  ENGLAND: United Kingdom
CUMENT TYPE:
                  Journal; Article; (JOURNAL ARTICLE)
                  English
ANGUAGE:
                  Priority Journals
LE SEGMENT:
TRY MONTH:
                  198406
ITRY DATE:
                  Entered STN: 19900319
                  Last Updated on STN: 19900319
                  Entered Medline: 19840608
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nurification* of both S-100

that is described for the re

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***protein*** and calmodular from crude bovine brain extracts by the use of a fluphenazine-Sepharose attricts column ***eluted*** states with decreasing concentrations of free Ca2+. ***Protein*** containing only
   alpha-subunit was ***purified*** from preparations of S-100
***protein*** by ***anion*** - ***exchange***
***chromatography*** . This ***protein*** co-migrated with the alpha-subunit of S-100 ***protein*** on sodium dodecyl sulphate/
     ***urea*** /polyacrylamide-gel electrophoresis and had an amino acid
   composition identical with that previously reported for this subunit. The
   results of u.v.-absorption and fluorescence-emission spectroscopy indicate
   that the tryptophan residue of the ***purified*** alpha-subunit of
           ***protein*** undergoes a Ca2+-induced change in environment.
   Measurements of changes in tryptophan fluorescence with increasing Ca2+
   concentrations suggest an apparent dissociation constant of the
   alpha-subunit for Ca2+ of 7 X 10(-5)M in the absence of K+. In the
   presence of 90mM-K+ this value is increased to 3.4 X 10(-4)M.
   ANSWER 32 OF 37 CAPLUS COPYRIGHT 2003 ACS
 ESSION NUMBER: 1984:47927 CAPLUS
 UMENT NUMBER:
                          100:47927
                         Ion-exchange chromatography of proteins using
'LE:
                        diethylaminoethyl-Sephadex in the presence of urea
HOR(S): Kadoya, Toshihiko; Okuyama, Tsuneo PORATE SOURCE: Fac. Sci., Tokyo Metrop. Univ., Tokyo, Japan
                          Bunseki Kagaku (1983), 32(11), 664-9
                          CODEN: BNSKAK; ISSN: 0525-1931
JUMENT TYPE:
                          Journal
   Japanese
To ***purify*** cytoskeletal ***proteins*** such as glial
fibrillar acidic ***protein*** (GFA) or neurofilament ***proteins***
(NFP), the ***proteins*** were extd. with 2M ***urea*** from brain
IGUAGE:
   tissue. ***Anion*** - ***exchange*** ***chromatog*** . was done
   on a 6 .times. 0.75 cm (inner diam.)-column packed with DEAE-Sephadex A 50
   equilibrated in 10 mM K phosphate buffer (pH 7.1) with or without
     ***urea*** , and a linear NaCl gradient was used for ***elution***
     ***proteins*** . Human IgG, bovine serum albumin (BSA), and human serum ***proteins*** were used as stds. In the presence of ***urea*** ,
   interaction between the ion-exchanger and BSA was reduced. In the
   presence of 6M ***urea*** , the ***elution*** patterns of the
     ***proteins*** were affected; however, 2M ***urea*** scarcely
   produced an effect on ion-exchange chromatog. of human serum
     ***proteins*** . A 2M ***urea*** ext. of bovine brain, which was
   rich in GFA and NFP, was sepd. on DEAE-Sephadex A 50 in the presence of 2M
     ***urea***
2 ANSWER 33 OF 37 CAPLUS COPYRIGHT 2003 ACS
                                                             DUPLICATE 23
CESSION NUMBER: 1982:595277 CAPLUS
CUMENT NUMBER:
                          97:195277
                          Purification of the glial fibrillary acidic protein by
TLE:
                          anion-exchange chromatography
                          Dahl, Doris; Crosby, Carol J.; Gardner, Eileen E.;
THOR(S):
                          Bignami, Amico
                          Spinal Cord Injury Res. Lab., West Roxbury Veterans
RPORATE SOURCE:
                          Adm. Med. Cent., Boston, MA, 02132, USA
                          Analytical Biochemistry (1982), 126(1), 165-9
URCE:
                          CODEN: ANBCA2; ISSN: 0003-2697
CUMENT TYPE:
                          Journal
                          English
NGUAGE:
   A procedure for the ***isolation*** of assembly-competent glial
   fibrillary acidic (GFA) ***protein*** from 2M ***urea*** exts. of bovine spinal cord by ***anion*** - ***exchange*** ***chromatog***
                                                                      ***chromatog***
   . is reported. The tissue was previously extd. with low-ionic-strength
   buffer. The procedure allowed the sepn. of nondegraded GFA
     ***protein*** from GFA ***protein*** comprising degraded species.
   As previously reported for neurofilament prepns. obtained from porcine
   spinal cord by N. Geisler and K. Weber (1981), the procedure also allowed
   the simultaneous sepn. of the 3 neurofilament ***polypeptides***
   (200,000; 150,000; and 70,000 daltons) contained in the 2M ***urea***
                                                                 ***eluted***
   ext. Brain filament ***proteins*** sequentially
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increasing salt concn. (25-200 mM NaCl) according to their isoelec. point.

Proteins with higher pI ***eluted*** first. Tubulin

polypeptides ANSWER 34 OF 37 CAPLUS COPYRIGHT 2003 ACS 1980:617475 CAPLUS ESSION NUMBER: CUMENT NUMBER: 93:217475 Radioligand assays - methods and applications. LE: 125I-monoiodoinsulin - preparation, immunological and biological characterization Besch, W.; Woltanski, K. P.; Knospe, S.; Ziegler, M.; THOR (S): Keilacker, H. Abt. Radioimmunol., Zentralinst. Diabetes "Gerhardt RPORATE SOURCE: Katsch", Karlsburg, 2201, Ger. Dem. Rep. Acta Biologica et Medica Germanica (1980), 39(4), JRCE: 495-502 CODEN: ABMGAJ; ISSN: 0001-5318 Journal CUMENT TYPE: **VGUAGE:** German Monoiodoinsulin was prepd. from a heterogeneous 125I-iodination mixt. by ***anion*** - ***exchange*** ***chromatog*** . on DEAE-Sephadex A 25 without gradient ***elution*** (40 mM Tris, 0.1M NaCl, and 7M ***urea*** at pH 9.0 and 4.degree.). The sp. radioactivity of [125I] monoiodoinsulin was 14.3 TBq/g, i.e., an I content of 1.04 atoms/mol. ***insulin*** . Monoiodoinsulin was indistinguishable from ***insulin*** with respect to binding to guinea pig antiinsulin ***isolated*** ***receptors*** of serum and to ***insulin*** rat adipocytes. The biol. potency (96.5% of the immunoreactive ***insulin*** activity) detd. by the conversion of [1-14C]D-glucose to 14CO2 in vitro by rat fat cells was similar to that of native ***insulin*** ANSWER 35 OF 37 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.DUPLICATE CESSION NUMBER: 1978:214900 BIOSIS CUMENT NUMBER: BA66:27397 HETEROGENEITY POLY PEPTIDE CHAIN COMPOSITION AND ANTIGENIC TLE: REACTIVITY OF COMPLEMENT C-3 NEPHRITIC FACTOR. DAHA M R; AUSTEN K F; FEARON D T THOR(S): DEP. MED., LEIDEN UNIV., LEIDEN, NETH. RPORATE SOURCE: J IMMUNOL, (1978) 120 (4), 1389-1394. URCE: CODEN: JOIMA3. ISSN: 0022-1767. LE SEGMENT: BA; OLD NGUAGE: English C3 [3rd component of complement] nephritic factor (C3NeF), recognized by its capacity to stabilize the cell-bound amplification C3 convertase, ***purified*** from sera of 3 patients with C3b, Bb, was hypocomplementemic glomerulonephritis and of 2 patients with partial lipodystrophy by QAE-A50 Sephadex and SP C-25 Sephadex chromatography, affinity for the fluid phase amplification C3 convertase and QAE-A50 Sephadex chromatography. Each C3NeF preparation exhibited heterogeneity during cation exchange chromatography. The isoelectric points of the ***eluted*** fractions ranged at pI 8.3-8.9. The chromatographic fractions were interacted with ***purified*** B, .hivin.D and C3 to form fluid phase C3b,Bb(C3NeF) which sedimented as a 10S complex on sucrose density gradient ultracentrifugation; the ***isolated*** convertase was decayed with release of C3NeF, which was separated from C3b ***anion*** ***exchange*** and Bi [inactivated Factor B] by ***Purified*** ***chromatography*** . preparations of C3NeF radiolabeled with 125I were bound from 92-98% by 109 erythrocytes bearing C3b, Bb. Erythrocytes carrying C3b bound from 0.6-18% and EA [antibody-sensitized erythrocytes] engaged in no specific uptake. Analysis of all 125I-C3NeF preparations by SDS-PAGE [sodium dodecyl sulfate-polyacrylamide gel electrophoresis] demonstrated an apparent MW of 150,000. After reduction in the presence of 8M ***urea*** , each ***polypeptide*** chains of 54,000 and 125I-C3NeF preparation revealed 23,500 MW which corresponded with the positions of the H and L chains of reduced Ig[immunoglobulin]G. The reaction of 125I-C3NeF from 4 patients was positive with Sepharose-bound antisera to IgG, .alpha.1, .alpha.2, .kappa. and .lambda. and negative with antisera to .mu., .alpha., .delta., .gamma.3 and .gamma.4. C3NeF from the 5th patient differed in not reacting

with antiserum to .kappa.. C3NeF is probably an autoantibody directed against the antigens expressed by the amplification C3 convertase, C3b,Bb.

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12 ANSWER 36 OF 37 CAPLUS COPYRECHT 2003 ACS
CCESSION NUMBER: 1976:16132
                                     CAPLUS
                       84:161321
OCUMENT NUMBER:
                       Purification of factor XII (Hageman factor) from human
ITLE:
                        plasma
                        Chan, John Y. C.; Movat, Henry Z.
JTHOR(S):
                       Inst. Immunol., Univ. Toronto, Toronto, ON, Can.
ORPORATE SOURCE:
                        Thrombosis Research (1976), 8(3), 337-49
OURCE:
                        CODEN: THBRAA; ISSN: 0049-3848
OCUMENT TYPE:
                        Journal
                        English
ANGUAGE:
   Blood-coagulation factor XII was ***purified*** from human plasma in 4
   steps in the presence of hexadimethrine bromide and soybean trypsin
   inhibitor (SBTI). After adsorption with aluminum hydroxide, the plasma was pptd. with polyethylene ***glycol*** . The ***protein***
   precipitating between 4.0 and 16% satn. was redissolved in .apprx.30% of
   the starting plasma vol. and chromatographed. The 1st chromatographic step was ***anion*** - ***exchange*** ***chromatog*** . on
   QAE-Sephadex, in the presence of hexadimethrine bromide and SBTI. The
   SBTI was used also during the 1st half of chromatog. on CM-Sephadex, from which factor XII ***eluted*** in the 2nd half. As a 4th preparative
   step factor XII was subjected to either gel filtration on Sephadex G 100
   or affinity chromatog. The latter consisted of an immunoadsorbent column,
   the antibody being against contaminating ***proteins***
                      from factor XII-deficient plasma. The final product
     ***isolated***
   exhibited a sharp intense band by Na dodecyl sulfate disc gel
   electrophoresis, with an estd. mol. wt. of 78,000.
12 ANSWER 37 OF 37 CAPLUS COPYRIGHT 2003 ACS
CCESSION NUMBER: 1971:135269 CAPLUS
OCUMENT NUMBER:
                        74:135269
                       Urate-binding alpha1-2 globulin. Isolation and
ITLE:
                      characterization of the protein from human plasma
                       Aakesson, I.; Alvsaker, J. O.
JTHOR(S):
                   Dep. Biochem., Univ. Oslo, Bindern, Norway
ORPORATE SOURCE:
OURCE:
                       European Journal of Clinical Investigation (1971),
                        1(4), 281-7
                        CODEN: EJCIB8; ISSN: 0014-2972
OCUMENT TYPE:
                        Journal
ANGUAGE:
                       English
   The urate-binding .alpha.1-2-globulin occurred mainly in the human plasma ***protein*** fraction ***eluted*** from DEAE-Sephadex columns with
   0.01M Na phosphate buffer, pH 7.35, contg. 0.40M NaCl. By
                                                                  ***anion***
      columns, followed by (NH4)2SO4 pptn. and preparative polyacrylamide gel
   electrophoresis, this urate-binding globulin was ***isolated***
   highly ***purified*** state. It was a rod-shaped glycoprotein with a
   mol. wt. of 67,000 as detd. by dodecyl sulfate-polyacrylamide gel
   electrophoresis and had an isoelec. point of pH 4.6. In the presence of
        ***urea*** and mercaptoethanol the ***protein*** did not split
   into subunits, indicating that it might represent a single
     ***polypeptide*** chain. The urate-binding globulin contained 12.1%
   carbohydrate, including galactose, mannose, galactosamine, and sialic
   acid. Its amino acid compn. did not differ significantly from that of
   other plasma ***proteins*** except for the presence of an unidentified
   compn. The urate-binding globulin did not seem to be identical to any
   previously characterized ***protein***
                                              from human blood.
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